What is claimed is:

- An isolated polynucleotide comprising a nucleotide sequence that is at least 80% identical to SEQ ID NO:1, and wherein a virus encoded by said polynucleotide infects mice.
- 2. The polynucleotide of claim 1 wherein said sequence is at least 95% identical to SEQ10 ID NO:1.
 - 3. The polynucleotide of claim 1, wherein said sequence is identical to SEQ ID NO:1.
 - 4. The polynucleotide of claim 1, wherein said polynucleotide consists of SEQ ID NO:1.
 - 5. An isolated polynucleotide comprising a sequence that is completely complementary to the sequence of claim 1.
 - 6. An amino acid sequence encoded by the nucleotide sequence of claim 1.
 - 7. A peptide of at least 20 amino acids encoded by the polynucleotide of claim 1.
 - 8. A peptide of at least 20 amino acids encoded by the polynucleotide of claim 2.
- 9. An isolated polypeptide comprising an amino acid sequence at least 80% identical to SEQ ID NO:2, and wherein said amino acid sequence is a murine norovirus sequence.
 - 10. The isolated polypeptide of claim 9 wherein the sequence is at least 95% identical to SEQ ID NO:2.

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- 11. The isolated polypeptide of claim 9 wherein the sequence is identical to SEQ ID NO:2.
- 12. An isolated polypeptide comprising an amino acid sequence at least 80% identical to
 SEQ ID NO:3, and wherein said amino acid sequence is a murine norovirus sequence.
 - 13. The isolated polypeptide of claim 12, wherein the sequence is at least 95% identical to SEQ ID NO:3.
- 10 14. The isolated polypeptide of claim 12 wherein the sequence consists of SEQ ID NO:3.
 - 15. An isolated polypeptide comprising an amino acid sequence at least 80% identical to SEQ ID NO:4, and wherein said amino acid sequence is a murine norovirus sequence.
- 16. The isolated polypeptide of claim 15 wherein the sequence is at least 95% identical to SEQ ID NO:4.
 - 17. The isolated polypeptide of claim 15 wherein the sequence consists of SEQ ID NO:4.
- 20 18. An isolated nucleotide sequence comprising at least 10 contiguous nucleotides completely complementary to the polynucleotide of claim 1, wherein said nucleotide sequence hybridizes to SEQ ID NO:1 under highly stringent conditions.
- 19. The isolated nucleotide sequence of claim 18 wherein said at least 10 contiguous25 nucleotides are completely complementary to SEQ ID NO:1.
 - 20. An isolated nucleotide sequence comprising at least 10 contiguous nucleotides identical to the polynucleotide of claim 1, wherein said nucleotide sequence hybridizes to the complement of SEQ ID NO:1 under highly stringent conditions.

- 21. The isolated nucleotide sequence of claim 20 wherein said at least 10 contiguous nucleotides are identical to SEQ ID NO:1.
- 22. A host cell transfected with the polynucleotide of claim 1, or a portion thereof that
 5 encodes a polypeptide of at least 20 amino acids.
 - 23. A vector comprising the polynucleotide of claim 1, or a portion thereof that encodes a polypeptide of at least 20 amino acids.
- 10 24. A host cell comprising the vector of claim 23.
 - 25. A method for detecting the presence of murine calicivirus in a sample, comprising
 - a) contacting the sample with the isolated nucleotide sequence of claim 18;
 - b) providing conditions that allow specific hybridization of the nucleotide sequence of step a) to the murine calicivirus; and
 - c) detecting whether the nucleotide sequence hybridizes to murine calicivirus.
 - 26. The method of claim 25 wherein said detecting step further comprises quantitating the presence of murine calicivirus.

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- 27. A method for detecting the presence of murine calicivirus in a sample, comprising
- a) contacting the sample with a nucleotide sequence capable of specifically hybridizing to murine calicivirus;
- b) providing conditions that allow specific hybridization of the nucleotide sequence of step a) to the murine calicivirus; and
 - c) detecting whether the nucleotide sequence hybridizes to murine calicivirus.
- 28. The method of claim 27 wherein said detecting step further comprises quantitating the presence of murine calicivirus.

- 29. An isolated polynucleotide encoding a peptide of a murine calicivirus, said polynucleotide consisting essentially of:
- a) a nucleotide sequence selected from the group consisting of nucleotides 147-5021 of SEQ ID NO:1; nucleotides 5057-6679 of SEQ ID NO:1; and nucleotides 6682-7302 of SEQ ID NO:1;
 - b) a nucleotide sequence that is at least 80% identical to the polynucleotide of a);
 - c) the complement of a) or b).
- 30. A vector comprising the polynucleotide of claim 29.

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- 31. A host cell transfected with the polynucleotide of claim 30.
- 32. An antibody capable of selectively binding to a murine calicivirus amino acid sequence, wherein said amino acid sequence is selected from the group consisting of SEQ
 ID NO:2, SEQ ID NO:3, SEQ ID NO:4, the amino acid sequence of the peptide of claim 7, and the amino acid sequence of the peptide of claim 8.
 - 33. The antibody of claim 32, wherein said antibody is a monoclonal antibody.
- 20 34. The antibody of claim 32, wherein said antibody is a polyclonal antibody.
 - 35. A method for identifying an agent capable of modulating or preventing murine calicivirus (MNV-1) infection, said method comprising
 - a) providing a mouse infected with MNV-1;
 - b) administering said agent to the mouse of a);
 - c) monitoring the outcome of the MNV-1 infection.